



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/015,989A

DATE: 09/17/2004

TIME: 13:35:44

Input Set : A:\ORT1552 Seq Listing 9-2004.txt
 Output Set: N:\CRF4\09172004\J015989A.raw

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3 <110> APPLICANT: Darrow, Andrew
4      Qi, Jenson
5      Andrade-Grodon, Patricia
7 <120> TITLE OF INVENTION: Zymogen Activation System
9 <130> FILE REFERENCE: ORT-1552
11 <140> CURRENT APPLICATION NUMBER: 10/015,989A
12 <141> CURRENT FILING DATE: 2001-12-10
14 <150> PRIOR APPLICATION NUMBER: 09/303,162
15 <151> PRIOR FILING DATE: 1999-04-30
17 <160> NUMBER OF SEQ ID NOS: 60
19 <170> SOFTWARE: PatentIn version 3.2
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22 <211> LENGTH: 361
23 <212> TYPE: DNA
24 <213> ORGANISM: Artificial
26 <220> FEATURE:
27 <223> OTHER INFORMATION: Fusion Gene Vectors
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34 gtggacgcgg ccgcctttgc tgccccctt gatgatgatg acaagatcg tgggggctat 180
36 gctctagata gcggcccgctt cccttagtg agggtaatg ctgcgacgac acatgataag 240
38 atacattgtat gagtttggac aaaccacaac tagaatgcag tgaaaaaaaaat gctttattt 300
40 taaaaatttgt gatgctatttgc ctttatttgt aaccattata agctgcaata aacaagtgt 360
42 c
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46 <211> LENGTH: 301
47 <212> TYPE: DNA
48 <213> ORGANISM: Artificial Sequence
50 <220> FEATURE:
51 <223> OTHER INFORMATION: Fusion Gene Vectors
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56 tttgatgatg atgacaagat cgttgggggc tattgtctag atacccctac gatgtgccc 120
58 attacgccta gcggcccgctt cccttagtg agggtaatg ctgcgacgac acatgataag 180
60 atacattgtat gagtttggac aaaccacaac tagaatgcag tgaaaaaaaaat gctttattt 240
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68 <211> LENGTH: 484
69 <212> TYPE: DNA
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:

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73 <223> OTHER INFORMATION: Fusion Gene Vector
 75 <400> SEQUENCE: 3

76	gaattcacca	ccatggacag	caaaggttcg	tcgcagaaaat	ccgcctgct	cctgctgctg	60
78	gtggtgtcaa	atctactctt	gtgccagggt	gtggctccg	actacaaggaa	cgacgacgac	120
80	gtggacgcgg	ccgctttgc	tgccttttt	atcgaggggc	gcattgtgga	gggctcgat	180
82	ctagataccc	ctacgtgt	cccgattacg	ccgctagata	cccctacat	gtgcccatt	240
84	acggccgtag	ataccactac	gatgtgccc	attacgcgc	tagataccccc	tacgtatgtc	300
86	ccgattacgc	ctagcggccg	cttccttta	gtgagggtta	atgcttcgag	cagacatgat	360
88	aagatacatt	gatgagtttgc	gacaaaccac	aactagaatg	cagtaaaaaa	aatgttttat	420
90	tttgaaattt	tgtgtatcta	ttgctttattt	tgttaaccatt	ataagctgca	ataaacaagt	480
92	tgac						484

95 <210> SEQ ID NO: 4
 96 <211> LENGTH: 382
 97 <212> TYPE: DNA

98 <213> ORGANISM: Artificial Sequence
 100 <220> FEATURE:
 101 <223> OTHER INFORMATION: Fusion Gene Vectors
 103 <400> SEQUENCE: 4

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106	gtggtgtcaa	atctactctt	gtgccagggt	gtggctccg	actacaaggaa	cgacgacgac	120
108	gtggacgcgg	ccgctttgc	tgccttttt	atcgaggggc	gcattgtgga	gggctcgat	180
110	aactgtctag	acatcaccat	caccatca	agcggccgct	tcccttagt	gagggttaat	240
112	gtctcgagca	gacatgataa	gatacattga	tgagtttgg	caaaccacaa	ctagaatgca	300
114	gtgaaaaaaa	tgctttattt	gtgaaatttgc	tgtatgttattt	gtcttatttgc	taaccattat	360
116	aagctgcaat	aaacaagttg	ac				382

119 <210> SEQ ID NO: 5
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 121 <212> TYPE: DNA

122 <213> ORGANISM: Artificial Sequence
 124 <220> FEATURE:
 125 <223> OTHER INFORMATION: Fusion Gene Vectors
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130	ttcggctcg	gggtccccga	ctacaaggac	gacgacgacg	cggccgcct	tgctgcccc	120
132	tttgatgt	atgacaagat	cgttggggc	tatgtcttag	acatcaccat	caccatca	180
134	agcggccgct	tcccttagt	gagggttaat	gttcgagca	gacatgataa	gatacattga	240
136	tgagtttgg	caaaccacaa	ctagaatgca	gtgaaaaaaa	tgcttatttgc	taaccattat	300
138	tgtatgttattt	gtcttatttgc	aaacaagttg	ac			352

141 <210> SEQ ID NO: 6
 142 <211> LENGTH: 385
 143 <212> TYPE: DNA

144 <213> ORGANISM: Artificial Sequence
 146 <220> FEATURE:
 147 <223> OTHER INFORMATION: Fusion Gene Vectors
 149 <400> SEQUENCE: 6

150	gaattcacca	ccatggctt	cctctggctc	ctctcctgct	gggcctcct	ggtaccacc	60
152	ttcggctcg	gggtccccga	ctacaaggac	gacgacgacg	cggccgcct	tgctgcccc	120
154	tttgatgt	atgacaagat	cgttggggc	tatgtcttag	ataccctac	gatgtgccc	180
156	attacgccc	tagacatcac	catcaccatc	actagcggcc	gttccctt	agtgagggtt	240

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PATENT APPLICATION: US/10/015,989A

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Output Set: N:\CRF4\09172004\J015989A.raw

158 aatgcttcga gcagacatga taagatacat ttagtggatggacaaacca caactagaat 300
 160 gcagtaaaaaa aaatgttttta tttgtgaaat ttgtgtatgc attgttttat ttgttaaccat 360
 162 tataagctgc aataaacaag ttgac 385
 165 <210> SEQ ID NO: 7
 166 <211> LENGTH: 1169
 167 <212> TYPE: DNA
 168 <213> ORGANISM: Artificial Sequence
 170 <220> FEATURE:
 171 <223> OTHER INFORMATION: Fusion Gene With Homo Sapien Serine Protease Catalytic Domain
 173 <400> SEQUENCE: 7
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 176 gtgggtgtcaa atctactctt gtgccagggt gtggctctcc actacaagga cgacgacgac 120
 178 gtggacgcgg cgccttgc tgccccctt gatgtatgt acaagatcg tgggggctat 180
 180 gctctagagg ccggcgtcg gcccggcag gtcagcatca cctatgaagg cgtccatgtg 240
 182 tgggtggct ctctcggtc tgagggtcg gtgtgtcg ctgctactg cttccccagc 300
 184 gagcaccaca aggaaggcta tgaggtaag ctggggggcc accagctaga ctcttactcc 360
 186 gaggacgcgc aggtcagcac cctgaaggac atcatcccc accccagcta cttccaggag 420
 188 ggctcccagg ggcacattgc actcctccaa ctcagcagac ccatcacctt ctccgcgtac 480
 190 atccggccca tctgcctccc tgagccaac gcctccttcc ccaacggcct ccactgcact 540
 192 gtcactggct ggggtcatgt ggccccctca tgagcctcc tgacgcccggccactgcag 600
 194 caactcgagg tgcctctgtat cagtcgtgag acgtgtact gcctgtacaa catcgacgc 660
 196 aaggctgagg agccgcactt tgtccaagag gacatgggtgt gtgtgtgcta tgggggggg 720
 198 ggcaaggacg cctggcagggt tgactctggg gcccactct cctggccctgt ggagggtctc 780
 200 tggtaacctga cgggcattgt gagctgggg gatgcctgtg gggccggccaa caggctgggt 840
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 204 ctcgtgtgg tgccccaaac ccaggagtc cagccgcaca gcaacctctg tggcagccac 960
 206 ctggccttca gctctagaca tcaccatcac catcaactgc ggccgcttcc ctttagtgag 1020
 208 ggttaatgtc tcgagcagac atgataagat acattgtga gtttggacaa accacaacta 1080
 210 gaatgcagtg aaaaaaatgc ttatgtgaaatgtga tgctattgtct ttatgttaaa 1140
 212 ccattataag ctgcaataaa caagttgac 1169
 215 <210> SEQ ID NO: 8
 216 <211> LENGTH: 1142
 217 <212> TYPE: DNA
 218 <213> ORGANISM: Artificial Sequence
 220 <220> FEATURE:
 221 <223> OTHER INFORMATION: Fusion Gene With Homo Sapien Serine Protease Catalytic Domain
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 224 gaattcacca ccatggctt ctcctggctc ctctcctgct gggccctcct gggtaccacc 60
 226 ttccggctgcg gggccccgaa ctacaaggac gacgacgacg cggccgcctc tgctgcccc 120
 228 ttgtatgtatg atgacaagat cttttggggc tatgtcttag agggccgtca gtggccctgg 180
 230 caggtcagca tcacccatga aggccgtccat gtgtgtgggt gctctctcg gtctgagcag 240
 232 tgggtgtgt cagtcgtca ctgttcccc agcgagcacc acaaggaagc ctatgaggc 300
 234 aagctggggg cccaccagct agactctac tccgaggac ccaagggtcag caccctgaag 360
 236 gacatcatcc cccacccag ctacccatccag gagggtccc agggcgcacat tgcaactcctc 420
 238 caactcagca gacccatcac cttctccgc tacatccgc ccatctgcct ccctgcagcc 480
 240 aacgcctcc tccccaaacgg cttccactgc actgtcaactg gctggggtca tggggcccc 540
 242 tcagtgagcc tcctgacgcc caagccactg cagcaactcg aggtgcctct gatcagtcgt 600
 244 gagacgtgtactgcctgtca caacatcgac gccaaggctg aggagccgca ctttgtccaa 660
 246 gaggacatgg tgggtgtgttgcctgtgg ctatgtggag gggggcaagg acgcctggca ggggtactct 720

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248	gggggcccac	tctcctgccc	tgtggagggt	ctctggtacc	tgacggcat	tgtgagctgg	780
250	ggagatgcct	gtggggcccg	caacaggcct	ggtgtgtaca	ctctggcctc	cagctatgcc	840
252	tcctggatcc	aaagcaagg	gt gacagaactc	cagctcgtg	ttggtgc	ccccaa	900
254	tcccagcccg	acagcaacct	ctgtggcagc	cacctggcct	tcagctctag	acatcaccat	960
256	caccatca	act gcccgt	tcccttagt	gagggttaat	gcttcgagca	gacatgataa	1020
258	gatacattga	tgagtttgg	caaaccacaa	ctagaatgca	gtgaaaaaaa	tgcttattt	1080
260	gtgaaatttg	tgatgtatt	gttttattt	taaccattat	aagctgcaat	aaacaagtt	1140
262	ac						1142

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266 <211> LENGTH: 1049

267 <212> TYPE: DNA

268 <213> ORGANISM: Artificial Sequence

270 <220> FEATURE:

271 <223> OTHER INFORMATION: Fusion Gene With Homo Sapien Serine Protease Catalytic

Domain

273 <400> SEQUENCE: 9

274	gaattcacca	ccatggacag	caaagg	ttcg	tcgcagaaat	cccg	cctgct	ctgt	60
276	gtgggtgtcaa	atctactctt	gtgccagg	gtgg	gtctccg	actaca	agg	cgac	120
278	gtggacgcgg	ccgcttgc	tgc	cccc	ctt	gatgat	gt	acaagat	180
280	aactgtctag	aa	cccc	ccatc	gcag	cc	ttgg	ccagg	240
282	ctctgtggcg	gtgtc	ttgt	agg	ggcaac	tgg	ttcc	ttt	300
284	ccgaaataca	cagtac	ggg	gagacc	ac	gc	tttt	aaaa	360
286	gaaatacctg	tgg	ttc	agg	tc	atcc	ccac	ac	420
288	cacaaccatg	atct	gtat	gt	tctt	caact	g	ctg	480
290	aagcccatca	gc	ctt	ggc	gaga	tca	tttt	tttt	540
292	tggggactg	tc	acc	agg	tcc	tgc	act	gt	600
294	aaaatctt	cc	caga	aa	gt	gt	gg	at	660
296	gtctgtgcag	gc	agc	agc	aa	gggg	ctg	ac	720
298	gtgtgtgatg	gt	gc	act	cc	gg	cat	caca	780
300	gacaaac	tc	at	tt	ttt	ttt	ttt	ttt	840
302	ggcagcaagg	g	ct	ct	g	ac	cc	ttt	900
304	ggttaatgt	t	cg	ac	at	gt	ttt	gg	960
306	gaatgcagt	aa	aaa	at	ttt	ttt	ttt	ttt	1020
308	ccattataag	ct	g	ca	at	aa	ttt	ttt	1049

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312 <211> LENGTH: 1052

313 <212> TYPE: DNA

314 <213> ORGANISM: Artificial Sequence

316 <220> FEATURE:

317 <223> OTHER INFORMATION: Fusion Gene With Homo Sapien Serine Protease Catalytic

Domain

319 <400> SEQUENCE: 10

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322	gtgggtgtcaa	atctactctt	gtgccagg	gtgg	gtctccg	actaca	agg	cgac	120
324	gtggacgcgg	ccgcttgc	tgc	cccc	ctt	gatgat	gt	acaagat	180
326	aactgtctag	aa	agg	ccatc	cc	ggc	cc	tttt	240
328	ctctgtgggg	cgac	ct	cc	ttt	gg	ctt	ttt	300
330	cccc	ctt	ttt	cc	ttt	ttt	ttt	ttt	360
332	cc	cc	cc	cc	cc	cc	cc	cc	420
334	gacc	acc	cc	cc	cc	cc	cc	cc	480
336	gtgc	gac	cc	cc	cc	cc	cc	cc	540

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Output Set: N:\CRF4\09172004\J015989A.raw

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/17/2004
PATENT APPLICATION: US/10/015,989A TIME: 13:35:45

Input Set : A:\ORT1552 Seq Listing 9-2004.txt
Output Set: N:\CRF4\09172004\J015989A.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1

VERIFICATION SUMMARY

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DATE: 09/17/2004

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